## **AMENDMENTS TO THE CLAIMS:**

Claim 12 has been cancelled. Claims 1, 7, 8, 15, 16, 17, 20, 26, 30, 33, 34, 35, 38, 44, 49, 52, and 53 have been amended. Claims 1, 7, 8, 12, 15-17, 19, 20, 26, 30, 33-35, 37, 38, 44, 49, 52, and 53 are pending. The following is the status of the claims of the above-captioned application, as amended.

- 1. (Currently Amended) A method of producing a heterologous biological substance polypeptide, comprising:
- (a) cultivating a mutant of a parent Aspergillus niger strain in a medium suitable for the production of the heterologous biological-substance polypeptide, wherein (i) the mutant strain comprises a first nucleotide sequence encoding the heterologous biological-substance polypeptide and one or more second nucleotide sequences comprising a modification of glaA, and at least one of the genes selected from the group consisting of asa, amyA, amyB, prtT, and oah, and (ii) the mutant strain is deficient in the production of glucoamylase, and at least-one enzyme-selected from the group consisting of acid stable alpha-amylase, neutral alpha-amylase A, and neutral alpha-amylase B, protease, and oxalic acid hydrolase compared to the parent Aspergillus niger strain when cultivated under identical conditions; and
- (b) recovering the heterologous biological substance polypeptide from the cultivation medium.
- 2. (Canceled).
- 3. (Canceled).
- 4. (Canceled).
- 5. (Canceled).
- 6. (Canceled).
- 7. (Currently Amended) The method of claim 1, wherein the biological substance heterologous polypeptide encoded by the first nucleotide sequence is a biopolymer an antibody, antigen, antimicrobial peptide, enzyme, growth factor, hormone, immunodilator,

neurotransmitter, receptor, reporter protein, structural protein, or transcription factor.

8.	(Currently Amended) The method of claim 7, wherein the biopelymer enzyme is eelected
<del>from</del>	the group consisting of a nucleic acid, polyamide, polyamine, polyal, polypoptide, and
polys	<del>paccharide</del> an oxidoreductase, transferase, hydrolase, lyase, isomerase, or ligase.

- 9. (Canceled).
- 10. (Canceled).
- 11. (Canceled).
- 12. (Cancelled).
- 13. (Canceled).
- 14. (Canceled).
- 15. (Currently Amended) The method of claim 1, wherein the mutant strain produces at least 25% less enzyme for each of glucoamylase, and one or more enzymes selected from the group consisting of acid stable alpha-amylase, neutral alpha-amylase A, and neutral alpha-amylase B, protease, and oxalic acid hydrolase compared to the parent *Aspergillus niger* strain when cultivated under identical conditions.
- 16. (Currently Amended) The method of claim 1, wherein the mutant strain is completely deficient in glucoamylase, and at least one enzyme selected from the group consisting of acid stable alpha-amylase, neutral alpha-amylase A, and neutral alpha-amylase B, protease, and oxalic acid hydrolase compared to the parent *Aspergillus niger* strain when cultivated under identical conditions.
- 17. (Currently Amended) The method of claim 1, wherein the mutant strain further comprises a modification of one or more genes which encode an enzyme having proteolytic activity.
- 18. (Canceled).

- 19. (Original) The method of claim 1, wherein the mutant strain further comprises a modification of one or more genes encoding an enzyme selected from the group consisting of a carbohydrase, carboxypeptidase, catalase, cellulase, chitinase, cutinase, cyclodextrin glycosyltransferase, deoxyribonuclease, esterase, galactosidase, beta-galactosidase, glucose oxidase, glucosidase, haloperoxidase, hemicellulase, invertase, isomerase, laccase, ligase, lipase, lyase, mannosidase, oxidase, pectinolytic enzyme, peroxidase, phytase, phenoloxidase, polyphenoloxidase, ribonuclease, transferase, alpha-1,6-transglucosidase, transglutaminase, and xylanase.
- 20. (Currently Amended) A mutant of a parent Aspergillus niger strain, comprising a first nucleotide sequence encoding a heterologous biological-substance polypeptide and one or more second nucleotide sequences comprising a modification of glaA, and at least one of the genes selected from the group-consisting of asa, amyA, amyB, prtT, and oah, wherein the mutant strain is deficient in glucoamylase, and at least one enzyme selected from the group consisting of acid stable alpha-amylase, neutral alpha-amylase A, and neutral alpha-amylase B, protease, and oxalic acid hydrolase compared to the parent Aspergillus niger strain when cultivated under identical conditions.
- 21. (Canceled).
- 22. (Canceled).
- 23. (Canceled).
- 24. (Canceled).
- 25. (Canceled)
- 26. (Currently Amended) The mutant strain of claim 20, wherein the biological substance heterologous polypeptide encoded by the first nucleotide sequence is a biopolymer an antibody, antigen, antimicrobial peptide, enzyme, growth factor, hormone, immunodilator, neurotransmitter, receptor, reporter protein, structural protein, or transcription factor.

- 27. (Canceled).
- 28. (Canceled).
- 29. (Canceled).
- 30. (Currently Amended) The mutant strain of claim 20 26, wherein the biological-substance enzyme encoded by the first nucleotide sequence is a metabolite an oxidoreductase, transferase, hydrolase, lyase, isomerase, or ligase.
- 31. (Canceled).
- 32. (Canceled).
- 33. (Currently Amended) The mutant strain of claim 20, which produces at least 25% less enzyme for each of glucoamylase and one-or more enzymes selected from the group consisting of acid stable alpha-amylase, neutral alpha-amylase A, and neutral alpha-amylase B, protease, and oxalic acid hydrolase compared to the parent *Aspergillus niger* strain when cultured under identical conditions.
- 34. (Currently Amended) The mutant strain of claim 20, which is completely deficient in glucoamylase and one or more enzymes selected from the group consisting of acid stable alpha-amylase, neutral alpha-amylase A, and neutral alpha-amylase B, protease, and oxalic acid hydrolase compared to the parent *Aspergillus niger* strain when cultured under identical conditions.
- 35. (Currently Amended) The mutant strain of claim 20, which further comprises a modification of one or more genes which encode an enzyme having proteolytic activity.
- 36. (Canceled).
- 37. (Original) The mutant strain of claim 20, which further comprises a modification of one or more genes encoding an enzyme selected from the group consisting of a carbohydrase, carboxypeptidase, catalase, cellulase, chitinase, cutinase, cyclodextrin glycosyltransferase,

deoxyribonuclease, esterase, galactosidase, beta-galactosidase, glucose oxidase, glucosidase, haloperoxidase, hemicellulase, invertase, isomerase, laccase, ligase, lipase, mannosidase. oxidase. pectinolytic enzyme, peroxidase. phytase, phenoloxidase. polyphenoloxidase, ribonuclease, transferase, alpha-1,6-transglucosidase, transglutaminase, and xvlanase.

- 38. (Currently Amended) A method for obtaining a mutant of a parent *Aspergillus niger* strain, comprising:
- (a) introducing into the parent Aspergillus niger strain a first nucleotide sequence encoding a heterologous biological-substance polypeptide and one or more second nucleotide sequences comprising a modification of glaA, and at least one of the genes selected from the group consisting of asa, amyA, amyB, prtT, and oah; and
- (b) identifying the mutant strain from step (a) comprising the modified nucleotide sequence, wherein the mutant strain is deficient in the production of glucoamylase, and at least one enzyme selected from the group consisting of acid stable alpha-amylase, neutral alpha-amylase A, and neutral alpha-amylase B, protease, and oxalic acid hydrolase compared to the parent Aspergillus niger strain when cultivated under identical conditions.
- 39. (Canceled).
- 40. (Canceled).
- 41. (Canceled).
- 42. (Canceled).
- 43. (Canceled).
- 44. (Currently Amended) The method of claim 38, wherein the biological substance heterologous polypeptide encoded by the first nucleotide sequence is a biopolymer an antibody, antigen, antimicrobial peptide, enzyme, growth factor, hormone, immunodilator, neurotransmitter, receptor, reporter protein, structural protein, or transcription factor.
- 45. (Canceled).

- 46. (Canceled).47. (Canceled).
- 48. (Canceled).
- 49. (Currently Amended) The method of claim 38 44, wherein the biological substance enzyme encoded by the first nucleotide sequence is a metabolite an oxidoreductase, transferase, hydrolase, lyase, isomerase, or ligase.
- 50. (Canceled).
- 51. (Canceled).
- 52. (Currently Amended) The method of claim 38, wherein the mutant strain produces at least 25% less enzyme for each of glucoamylase, and one or more enzymes selected from the group consisting of acid stable alpha-amylase, neutral alpha-amylase A, and neutral alpha-amylase B, protease, and oxalic acid hydrolase compared to the parent *Aspergillus niger* strain when cultured under identical conditions.
- 53. (Currently Amended) The method of claim 38, wherein the mutant strain is completely deficient in glucoamylase, and one or more enzymes selected from the group consisting of acid stable alpha-amylase, neutral alpha-amylase A, and neutral alpha-amylase B, protease, and oxalic acid hydrolase compared to the parent *Aspergillus niger* strain when cultured under identical conditions.
- 54. (Canceled).
- 55. (Canceled).
- 56. (Canceled).